



Redesigning of Microbial Cell Surface and Its Application to Whole-Cell Biocatalysis and Biosensors

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Abstract Microbial cell surface display technology can redesign cell surfaces with functional proteins and peptides to endow cells some unique features. Foreign peptides or proteins are transported out of cells and immobilized on cell surface by fusing with anchoring proteins, which is an effective solution to avoid substance transfer limitation, enzyme purification, and enzyme instability. As the most frequently used prokaryotic and eukaryotic protein surface display system, bacterial and yeast surface display systems have been widely applied in vaccine, biocatalysis, biosensor, bioadsorption, and poly-peptide library screening. In this review of bacterial and yeast surface display systems, different cell surface display mechanisms and their applications in biocatalysis as well as biosensors are described with their strengths and shortcomings. In addition to single enzyme display systems, multi-enzyme co-display systems are presented here. Finally, future developments based on our and other previous reports are discussed.

Keywords Microbial cell surface display · Bacterial surface display · Yeast surface display · Carrier protein · Host cell · Passenger protein · Whole-cell biocatalysis · Biosensor · Multi-enzyme co-display system

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Introduction

Microbial surface display technology refers to the use of gene engineering means to realize presentation of foreign peptides or proteins on the surfaces of microorganisms in the forms of fusion proteins, which can maintain their relatively independent spatial structure and biological activity [1]. As the first microbial surface display system, the phage display system was established in the mid-80s of the twentieth century, where peptides and small proteins were displayed on the phage surface by fusing with the capsid protein of filamentous phage [2]. The phage display system has many remarkable advantages, such as high-throughput biopanning, screening of mimic epitopes, simple preparation process, and so on. In recent years, this technology has had a far-reaching influence on protein molecule mutual recognition, the development of new vaccines and tumor treatments [3]. With the development of surface display technology, surface display system has been expanded from phage display system to other microbial cell display systems [4]. Currently, bacterial and yeast surface display system is the most frequently used prokaryotic and eukaryotic protein surface display system, respectively, due to unique features, including diversity of carriers, high yields of recombinant proteins, and amenability to gene engineering [5].

A cell surface display system contains three factors, namely host, carrier, and passenger. The host cells serve as the matrix to bind fusion proteins of foreign protein and an anchoring motif. The carriers are some outer membrane protein and appendant of cell surface, whose signal peptides can facilitate the transverse of passengers from intracellular to surface. The passengers are the target foreign protein, which can be displayed on the surface of cell through this technology. Microbial cell surface display technology involves the process of membrane transportation, which is closely related to the mechanism of protein secretion. There are various protein display mechanisms in different cell types (Fig. 1). Therefore, it is indispensable to coordinate with the three essential elements to establish a successful surface display system.

Compared with free enzyme, cell surface-displayed enzyme has several advantages. Firstly, for the biotransformation of multi-step reaction, whole-cell catalysis can improve the efficiency of biocatalysis by coupling the cofactor and other enzymes in the cell. Secondly, the preparation of enzyme is simply achieved by culturing microbial cells, thus greatly simplifying the preparation process of the protein and reducing the production cost. Thirdly, the stability of cell surface-displayed enzyme is improved, and the whole-cell catalyst could be reusable without compromising its activity [1].

Until now, microbial surface display technology has been widely applied in all aspects of the biotechnology, such as live vaccine development, peptide library construction and screening, bioremediation of heavy metal pollution, whole-cell catalysis, and biosensors [4, 9].

Up to now, there have been several reviews related to recent advances on different surface display systems developed for phage, bacteria, and yeast [1, 10], as well as their applications in screening of protein libraries [6], vaccines [11], and medical and environmental aspects [12]. In this review, we will focus on bacterial and yeast surface display systems, as well as their applications on whole-cell biocatalysis and biosensors. Especially, constructions and performances of multi-enzyme co-display systems will be discussed.

Gram-Negative Bacterial Cell Surface Display

The cell envelope structure of Gram-negative bacteria is complex, composed of cytoplasmic membrane, periplasm, and outer membrane. In order to display on the outer membrane of cells, the foreign protein fused with the anchoring motif should pass through the cytoplasmic membrane

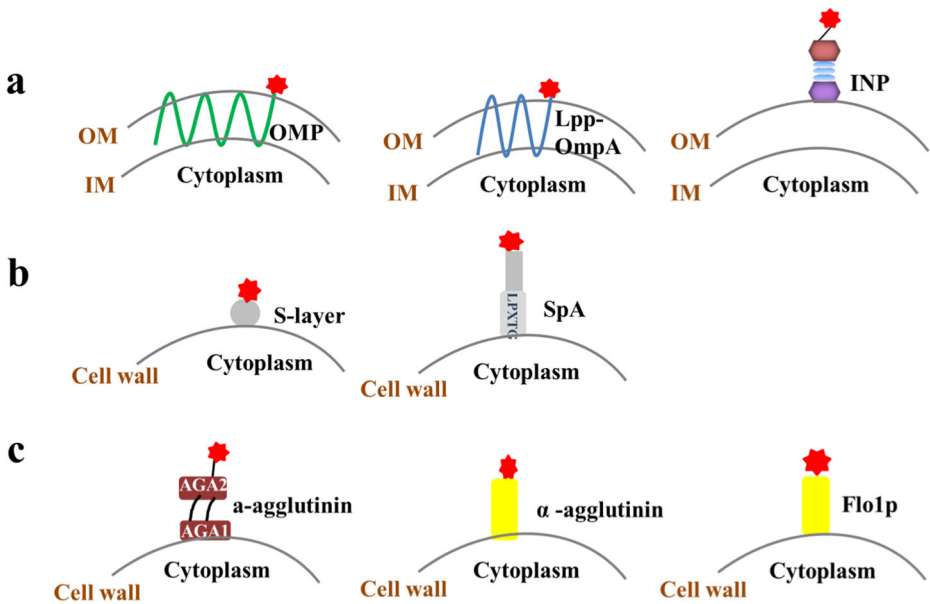


Fig. 1 Some typical carriers used in Gram-negative bacterial (a), Gram-positive bacterial (b), and yeast (c) cell surface display systems. Red stars represent passengers. OM refers to outer membrane. IM refers to inner membrane. OMP refers to outer membrane protein. Lpp-OmpA chimera comprises lipoprotein and outer membrane protein. INP refers to ice nucleation protein. SpA protein refers to *Staphylococcus aureus* protein A. Surface layers (S-layers) are outside of the cell structure of many Archaea and Bacteria. α - and a-agglutinin can mediate the mating between yeast cells. Flocculin Flo1p is flocculating protein of *Saccharomyces cerevisiae* (Ref. [6–8])

and periplasm space. Various display systems have been developed successfully, exhibiting different presentation profiles with diverse targeting and anchoring mechanisms of carrier proteins. The Gram-negative bacterium *Escherichia coli* is the most frequently used host in view of its mature genetic manipulation system and high yields of biomass. Representative display systems for Gram-negative bacterial surface display systems are shown in Table 1.

Outer Membrane Protein Display System

Outer membrane proteins (OMPs) span the outer membrane through antiparallel β -sheets which form a certain 3-D structure like a barrel, such as LamB (maltoporin), OmpA (out membrane protein A), and PhoE (phosphate-inducible porin) [39, 40]. The external loops can be inserted into foreign peptide fragments. Therefore, heterologous proteins can be fused with these loops for cell surface display. Most OMP mediate display systems are limited to small foreign protein display without the disruption of membrane integrity. Fortunately, a few of systems have been developed recently to facilitate stable display of large passengers. For example, amylase (77 KDa) could be localized on the surface of *E. coli* using protein PgsA from *Bacillus subtilis* as an anchoring motif [15].

Lipoprotein Display System

Lipoprotein with the lipid-modified cysteine of N-terminus was located at the periplasmic space. TraT has been used to display foreign peptides at two fusion sites, including the middle

Table 1 Representative display systems and applications for Gram-negative bacterium surface display systems

Carrier		Passenger	Application	Reference
OMPs	LamB	C3 and preS2 epitope	Vaccines	[13]
	PhoE	Antigenic determinants	Vaccines	[14]
	PgsA	Amylase	Biocatalysis	[15]
INP	N- and C-terminal	OPH	Screening of OPH variants	[16]
	N- and C-terminal	Alditol oxidase	Bioadsorption	[17]
	N- and C-terminal	Transglucosidase	Biocatalysis	[18]
	N-terminal	Xylose dehydrogenase	Biosensor	[19]
	N-terminal	Glucose dehydrogenase	Biosensor	[20]
	N-terminal	Triphenylmethane reductase	Biotransformation	[21]
	N-terminal	Green fluorescent protein	Report gene	[22]
	N-terminal	Laccase	Biocatalysis	[23]
	N-terminal	OPH	Biosensor	[24]
	N-terminal	Glucose oxidase	Biocatalysis	[25]
	N-terminal	Glutamate dehydrogenase	Biosensor	[26, 27]
	N-terminal	Phosphate-binding protein	Bioremediation	[28]
	N- and C-terminal	Hepatitis C virus core protein surface antigen	Vaccines	[29]
Lipoprotein	Lpp-OmpA	β -lactamase	Bioadsorption and vaccines	[30]
	TraT	Pre-S antigen	Vaccines	[31]
ATs	AIDA-I	Cholera toxin B subunit	Vaccines	[32]
	AIDA-I	Lipase	Biocatalysis	[33]
	AIDA-I	Nitrilase	Biocatalysis	[34]
	EstA	Lipase	Biocatalysis	[35]
	EstA	Lipase-specific chaperone	Screening of foldase variants	[36]
	EspP	Single-chain antibody fragment	Screening of binding motifs	[37]
	IgAP	Metallothionein	Bioremediation	[38]

and the C-terminus of TraT [12, 41]. The Lpp-OmpA chimera comprises the signal peptide and nine N-terminal residues of lipoprotein, as well as the residues 49-159 of OmpA [42]. The hybrid display system has been used to display several passengers, such as esterases, organophosphate hydrolases (OPH), and β -lactamase [43]. However, cell viability and enzyme activity could be influenced [44].

Ice Nucleation Protein Display System

Ice nucleation protein (INP) is an outer membrane protein in ice nucleation active bacterium, which can accelerate the formation of ice crystals in supercooled water [45]. It is involved with the genus *Pseudomonas* [46, 47], *Erwinia* [48], and *Xanthomonas* [49]. INP consists of three distinct domains, namely, N-terminal, C-terminal, and central repeating domain. The amino acids of the N-terminal domain are relatively hydrophobic and link the protein to the outer membrane through a glycosylphosphatidylinositol anchor, which is endowed with a particular desirable ability to display proteins (Fig. 2). Compared with other anchoring units, INP has several distinct characteristics due to its special structural and functional features: (i) It has been proved that effective bacterial surface display of foreign proteins can be achieved using INP as an anchoring motif with the full-length or truncated sequence, indicating that C-terminal and central repeating domains are not necessary for the anchoring function. Thus, surface display of passengers with large size could be realized when only N-terminal domain was served as the anchoring motif. (ii) Foreign proteins can be stably expressed in the host without degradation by intracellular or extracellular proteases. In contrast with Lpp-OmpA

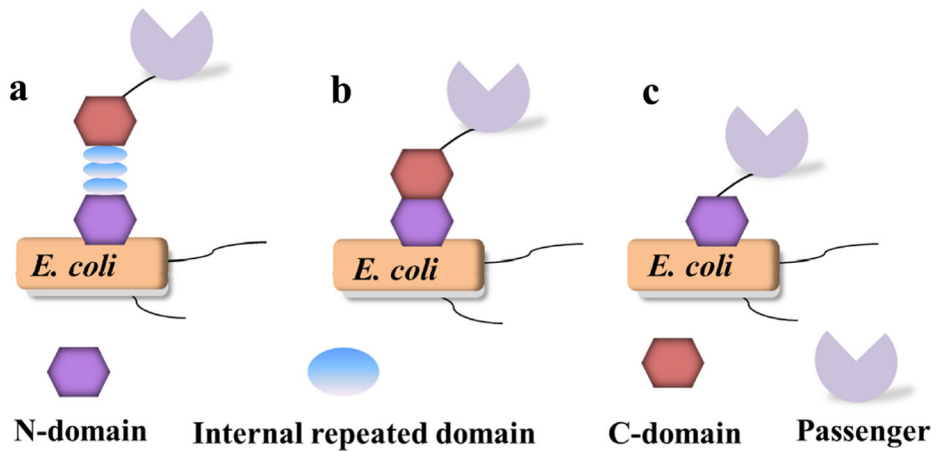


Fig. 2 Cell surface display systems using ice nucleation protein (INP) as carrier. **a** Display of passenger with the internal repeating domain, N domain, and C domain of INP. **b** Display of passenger with C domain and N domain of INP. **c** Display of passenger with only N domain of INP (Ref. [6])

chimera, the cell viability will not be disrupted when INP is displayed on the surfaces of bacteria [50]. (iii) INP has secreted, guided, and anchoring functions, which could make fusion proteins self-anchored on cell surface without requiring other specific accessory protein.

In recently years, INP has been widely used for surface display of heterologous proteins, even large proteins, such as Chitinase 92 (90 kDa) and cytochrome P450 (119 kDa) [51], or the combination of two proteins, such as organophosphate hydrolases (OPH) and methyl parathion (MPH) [52–54]. In the light of these results, the successful construction of INP-mediated bacterial surface display system and its application in biocatalysis and biosensor has been implemented in our laboratory. In detail, a variety of oxidoreductases and hydrolases were displayed on the surface of *E. coli* using N-terminal of INP as the anchoring motif. The excellent activity and stability of prepared whole-cell biocatalysts well exhibited the remarkable performance of INP display system. Novel biosensors were constructed by using the above whole-cell biocatalysts as detection elements, realizing detection quantification of the sugar and pesticides with high sensitivity and specificity [20, 24, 26, 55, 56]. However, this technique has some limitations. For example, for passenger proteins that contain disulfide bonds in the molecule, carrier proteins can interfere with the correct folding of the passenger protein, thereby reducing the activity of the passenger.

Autotransporter Protein Display System

Autotransporter proteins (ATs) are related to virulence of Gram-negative bacteria, and their main functions are to adhesion and protein hydrolysis (Table 2). ATs were firstly discovered in *Neisseria* sp., namely IgA protease [65]. All ATs have similar structural features. The N-terminal has a signal peptide that interacts with the Sec translocon. The central is the passenger region that carries the physiological translocation function. The C-terminal is called translocation unit (TU) [66]. Most of the signal peptide is composed of three parts, charged N region, hydrophobic H region and C region, and a signal peptidase recognition site. Sec translocon is able to recognize the N and H regions of the signal peptide to direct the polypeptide chain through the inner membrane into the periplasmic space, and then signal peptide is hydrolyzed and removed under the catalysis of the signal peptidase. The

Table 2 The types and functions of ATs

Type	Function	Source	Reference
Adhesin involved in diffuse adherence I (AIDA-I)	Adhesin	<i>Escherichia coli</i>	[57]
Extracellular serine protease (EspP)	Protease		[58]
Pertactin (Prn)	Adhesin and serum resistance	<i>Bordetella pertussis</i>	[59]
Immunoglobulin A1 protease (IgAP)	Protease and immunomodulatory	<i>Neisseria meningitidis</i>	[60]
Haemophilus adhesive protein (Hap)	Adhesin	<i>Haemophilus influenzae</i>	[61]
Esterase A (EstA)	Esterase	<i>Pseudomonas aeruginosa</i>	[62]
Moraxella catarrhalis adhesin (McaP)	Adhesin and protease	<i>Moraxella catarrhalis</i>	[63]
Intracellular spread autotransporter (IcsA)	Esterase	<i>Shigella flexneri</i>	[64]
Phospholipase B (PLB)	Phospholipase and adhesin	<i>Moraxella bovis</i>	[63]

size and function of the passenger region are very different, and the passenger area affects and determines the biological function of ATs. More than 97% of ATs are structurally right-handed beta-helices, which will correctly fold on cell surface after crossing the membrane, while the passenger region is away from the bacterial surface. The function of TU is to transport the passenger from the periplasmic space to the bacterial surface. At present, the 3-D structures of various TU have been reported, such as extracellular serine protease (EspP), adhesion involved in diffuse adherence I (AIDA-I) and EstA from *P. aeruginosa*, and so on. TU structure is basically similar, consisting of a β -barrel structure and an extended alpha-helix. The β -barrel structure consists of 12 consecutive β -sheets, and the α -helix is located in the inner cavity of the β -barrel structure. The adjacent two beta-sheets are linked together by a β -Turn or a cell exposed to an extracellular loop. Extracellular loop, β -turn, and α -helix can prevent the loss of intracellular material through the channel, but also to prevent the extracellular harmful substances (such as antibiotics, etc.) through the channel into the cell.

All the ATs use V-type secretory pathway. The secretory process consisted of four steps, including inner membrane transport, periplasmic transport, outer membrane transport, and passenger region modification [66]. The specific cell surface display mechanism using ATs as carrier was shown in Fig. 3. Many studies showed that AT was an ideal tool to display the exogenous proteins due to the fact that this system does not require the participation of other proteins and can transport larger size proteins as well as maintain their antigenic and biological functions [67]. For example, the outer membrane esterase EstA proved to successfully direct lipase and its specific chaperone to the microbial cell surface [36]. Besides, AT display system is highly efficient. The study has shown that ATs can exhibit more than 250,000 passenger protein on a single bacterial surface [34]. Coupling with flow cytometric analysis, the AT-mediated surface display system can be used to screen large libraries of variants generated by directed evolution. Although some models have been put forward to explain transport process of passenger region from periplasm to outer membrane, including the threading model, the multimeric model, the hairpin model, and so on, none of the above models can explain comprehensively the whole process, and part of the assumption is also the lack of experimental data support.

Gram-Positive Bacterial Cell Surface Display

Although Gram-positive bacteria have thicker and tougher outer cell wall than Gram-negative bacteria, proteins are presented on the surfaces of Gram-positive bacteria only through a single

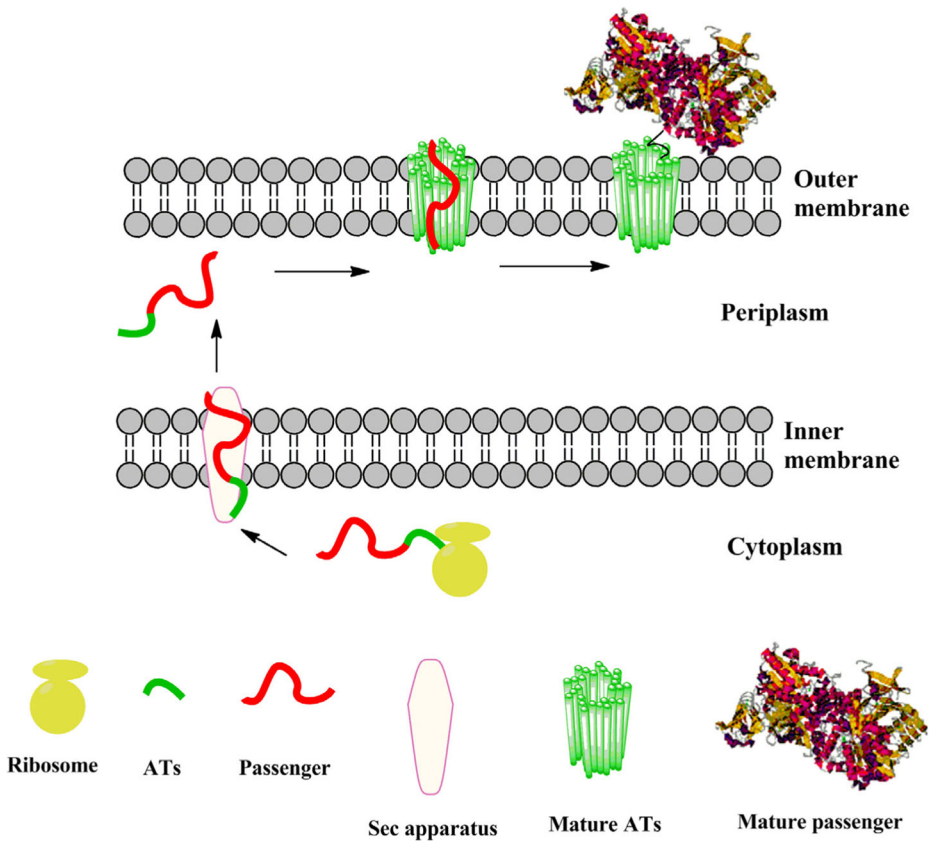


Fig. 3 The specific cell surface display mechanism using ATs as carrier. Sec translocon can recognize the signal peptide to direct the polypeptide chain through the inner membrane into the periplasmic space, and then signal peptide is hydrolyzed and removed under the catalysis of the signal peptidase. TU is folded into a β -barrel structure, which is then inserted into the extracellular membrane to form a channel through which the passenger is transported to the bacterial surface (Ref. [66])

membrane layer, avoiding negative influences on the structure and activity of protein during the traverse through the periplasmic space and the integration on the outer membrane. However, low transformation efficiency will hinder the construction of the peptide library with large capacity.

Staphylococcus aureus Protein A Display System

The C-terminal sequence of SpA can be anchored to the bacterial surface through the combination of a hydrophobic region with the peptidoglycan of cell wall [68]. The N-terminal is allowed to be inserted or replaced by foreign protein to achieve surface display. SpA consists of a signal peptide, an IgG-binding domain, and a C-terminal surface-binding domain. The C-terminal region consists of a charged repeating region containing LPXTG motif, a hydrophobic region, and a short charged tail RREL [69]. Staphylococcal surface display system utilizes the C-terminal anchoring region of SpA, replacing the IgG-binding domain with an exogenous protein, which is expressed as a fusion protein and displayed on the cell surface [70].

Surface Layer Display System

Surface Layers (S-layers) are outside of the cell structure of many Archaea and Bacteria, which is composed of a single protein or glycoprotein subunit and arranged regularly in the cell membrane or cell wall [71]. Amino acid sequence analysis showed that the S-layers of *Lactobacillus acidophilus* contain two conserved regions, including N-terminal signal sequence and C-terminal region, which can make foreign protein anchor on the surface of cell [72]. S-layers have been widely used in the construction of highly efficient secretory expression vector owing to their high level of transcription, expression, and secretion. Therefore, lactic acid bacterium surface display system was established using S-layers of *Lactobacillus* as carrier to display foreign antigens outside the cell surface, in order to avoid degradation by cytoplasmic protease [73].

Bacillus subtilis Spore Surface Display System

Bacillus subtilis spore surface display system is an important Gram-positive bacterial cell surface display system with spore coat protein as carrier, including CotB, CotC, CotG, OxdD, and other capsid proteins (Table 3). Studies showed that different carriers would lead to different display efficiencies [77]. The fusion of certain foreign proteins with the anchoring protein CotB does not affect the surface display [83]. But for some foreign proteins, the N-terminal fusion may be more reasonable. Isticato et al. displayed foreign protein using N-terminal and C-terminal of CotC protein. The result showed N-terminal fusion method could detect CotC monomer and dimer, but C-terminal fusion method could only detect a small amount of CotC monomer. Comparing with C-terminal fusion method, the display efficiency of N-terminal fusion method increased five times. The reason for this phenomenon may be that the C-terminal of CotC needs to interact with other CotC or capsid proteins during spore formation [84].

Yeast Cell Surface Display

Bacterial surface display system is incapable of expressing complex eukaryotic proteins, which need post-translational modifications to exhibit activity, including glycosylation and disulfide isomerization. Instead, as a eukaryotic system, *Saccharomyces cerevisiae* had been successfully employed to express and display dozens of complex proteins in the past decades, which could offer an easy handling procedure, stable activity of expressed enzymes, and feasibility to construct large protein libraries. In general, the target proteins were expressed in a fusion form

Table 3 *Bacillus subtilis* spore surface display systems

Carrier	Passenger	Application	Reference
OxdD/CotG	β -Glu/Phy	Biotransformation	[74]
CotB/CotC/CotG	UreA	Biotransformation	[75]
CotC	Ethanol dehydrogenase	Biocatalysis	[76]
CotB/CotC/CotG CotZ	FilD protein	Vaccines	[77]
CotC	Vp26	Vaccines	[78]
CotC	Human growth hormone	Vaccines	[79]
CotB/CotC	α -amylase	Biocatalysis	[80]
CotB	Tm1350/DSM	Biocatalysis	[81, 82]

with cell wall proteins, which were secreted out of the cell and then interact with the cell wall via a covalent or non-covalent linkage.

S. cerevisiae carrier protein can be divided into C-terminal anchored carrier protein and N-terminal anchored carrier protein according to its binding to cell wall (Table 4). C-terminal anchored carrier protein is mainly glycosylphosphatidylinositol (GPI) protein, such as α -agglutinin [95], Cwp1p [96], Cwp2p, and C-terminal of Flo1p [90], in which the N-terminal fragment was fused to passenger protein and C-terminal GPI anchor could covalently link to the cell wall through β -1,6-glucan bridge (Fig. 4a, b). N-terminal anchored carrier protein has three types, including a-agglutinin [97], N-terminal of Flo1p [98], and Pir protein [99]. The a-agglutinin is composed of two domains, namely Aga1 and Aga2. Aga1 could covalently link to the cell wall through GPI anchor. Aga2 binds to Aga1 by the disulfide bond, and the passenger could be display on cell surface by fusing with the C-terminal of Aga2 (Fig. 4c). But the display efficiency depends on the expression level of Aga1. The N-terminal of Flo1p is a flocculating domain that recognizes and is non-covalently bound to the mannan component in the cell wall, causing the cells to accumulate (Fig. 4d). Pir alternatively interacts with β -1,3-glucan in the cell wall via a ester bond. However, there is a lack of literature to show the display efficiency of these anchors which may vary dependent on the target proteins.

Applications of Microbial Surface Display in Biocatalysis

Biocatalyst generally refers to free or immobilized enzymes and living cells, which could be used to catalyze certain reaction with high efficiency under normal temperature and pressure. Microbial surface display system facilitates the development of new biocatalyst. Various enzymes have been presented on the surfaces of cells with favorable catalytic activity and stability.

Construction of Biocatalyst for Detoxification of Organophosphates

OPH was firstly displayed on the surface of *E. coli* JM105 by Lpp-OmpA display system, and the results showed that catalytic efficiency of whole cell was seven times higher than that of free enzyme towards parathion and paraoxon [100]. Unfortunately, the growth of cells could be inhibited grossly. Then, researchers targeted OPH on the surface of *E. coli* and *Pseudomonas putida* using truncated INP as anchoring unit in order to overcome this drawback. The results showed that enzyme activity of OPH displayed on the surface of *P. putida* was ten times higher

Table 4 Yeast surface display systems and their applications

Carrier	Passenger	Application	Reference
α -agglutinin	Endoglucanase	Biocatalysis	[85]
	Random protein	Screening of molecule	[86]
	Yeast metallothionein	Bioadsorbent	[87]
a-agglutinin	Glucanotransferase	Biocatalysis	[88]
	Integrin I domains	Protein evolution	[89]
C-terminal of Flo1p	Glucoamylase	Biocatalysis	[90]
N-terminal of Flo1p	Lipase	Biocatalysis	[91]
Pir1	Fucosyltransferase	Biocatalysis	[92]
Pir2	Mannosyltransferase	Biocatalysis	[93]
Pir4	Xylanase A	Biocatalysis	[94]

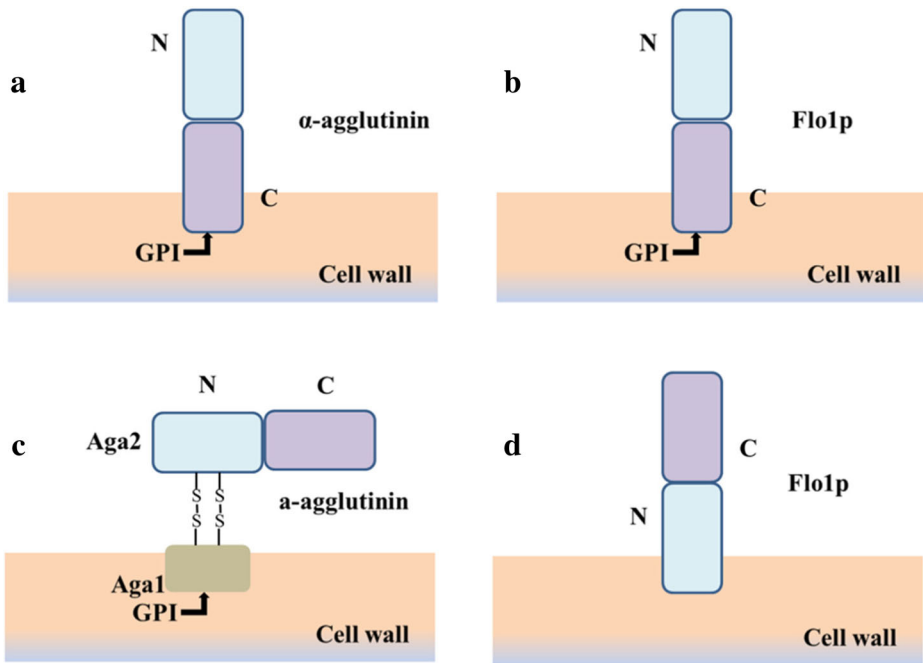


Fig. 4 Yeast surface display systems, including α -agglutinin (a), C-terminal of Flo1p (b), a-agglutinin (c), and N-terminal of Flo1p (d). α -agglutinin and C-terminal of Flo1p are C-terminal anchored carrier proteins, in which the N-terminal fragments were fused to passenger proteins and C-terminal GPI anchor could covalently link to the cell wall through β -1,6-glucan bridge. a-agglutinin and N-terminal of Flo1p are N-terminal anchored carrier proteins, in which the C-terminal fragments were fused to passenger proteins (Ref. [8])

than that on *E. coli* cell surface, probably because of the similar secretory channel between OPH original strain *Flavobacterium* sp. ATCC27551 and *P. diminuta* MG [44, 101]. *S. cerevisiae* also has been used as the host for surface display of OPH in consideration of practical industrial application, which exhibited about sixfold higher activity than INP display system [102]. In our previous work [24], OPH mutant S5 with improved enzyme activity was successfully displayed on the surface of *E. coli* using INP as anchoring motif and its level of enzyme activity (12.44 U/mg cells) was greatly higher than those of other reports (< 1 U/mg cells). The whole-cell catalyst was a multifunctional bacterium that could be used for degradation and detection of organophosphate pesticides. In subsequent work, at the aim of enhancing the activity and stability of OPH-fused whole-cell biocatalysts, the novel cell-inorganic hybrid materials were explored by the combination of microbial cell surface display technology and biomineralization (Fig. 5). Cell-displayed OPH showed the allosteric effect from “inactive” form to “active” form and the “active” form would be “fixed” when cell-displayed OPH was embedded into cobalt phosphate crystals. Therefore, the activity of mineralized OPH-fused cells was significantly enhanced to three times higher than that of original OPH-fused cells. As a bonus, the stability of the novel hybrid catalysts was also significantly improved. Further, the as-synthesized bio-inorganic hybrid catalysts could be applied on high-efficiency degradation and sensitive detection of paraoxon. To our knowledge, this work is the first example on the cell surface-displayed enzyme-inorganic hybrid biocatalysts and would provide a model method to develop a wide range of whole-cell biocatalysts for the diverse applications on industrial catalysis, environmental governance, and biosensing [103].

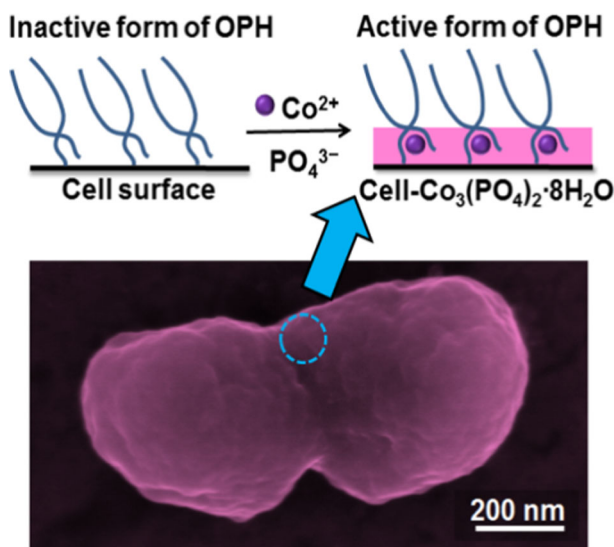


Fig. 5 The bio-inorganic hybrid whole-cell catalysts are fabricated for sensitive biosensing of organophosphorus pesticides by combination of biomineralization and microbial cell surface display technology. Due to the allosteric effect of organophosphorus hydrolase (OPH) embedded into inorganic crystal ($\text{Co}_3(\text{PO}_4)_2 \cdot 8\text{H}_2\text{O}$) from “inactive” form to “active” form, the biomineralized OPH-fused cells show the enhanced catalytic activity

Cell Surface Display of Lipase for Whole-Cell Biocatalysis

Since 1994, lipase has been displayed on the surfaces of cells based on various display systems. The common host and carrier were *E. coli* and OMPs, including OmpC, FadL, and OprF [104]. Activity of whole cell reached 2800 U/mg dry cells when lipase from *Bacillus* sp. strain TG43 presented on the surface of bacterium using OprF as carrier [105]. Lipase yeast display focused on α -agglutinin and Flo1p display system. *Rhizopus oryzae* lipase was first presented on the surface of yeast using N-terminal of Flo1p, which showed 14-fold higher activity (61.3 U/g dry cell weight) than α -agglutinin (4.1 U/g dry cell weight) [91]. Becker has found that the display efficiency of lipase and cell survival rate would decrease with the increasing molecular weight of target protein [106]. Compared with free lipase, the cell surface-displayed lipase showed superior stability towards high temperature and organic solvent [107]. Recently, new anchor proteins were developed, which included Flo9 and Pir1 from *Pichia pastoris*, and were able to display lipase B on yeast surface for hydrolysis of tributyrin. The constructed whole-cell biocatalyst showed conservation of $\sim 40\%$ of its original activity after 3 h incubation at 45 °C, while the free enzyme lost 60% of its original activity after 10 min at 45 °C [108]. Until now, display of lipase has been mainly applied to enantioselective resolution of racemic compounds. In addition, biodiesel could be catalyzed using cell surface-displayed lipase as whole cell biocatalyst [91, 109].

Construction of Multi-Enzyme Co-displayed Systems for Bioethanol Production

Compared to some yeast species such as *P. pastoris* and *Hansenula polymorpha*, *S. cerevisiae* has superior fermentation characteristics, especially high cell density culture on an economical carbon source. These features permit scientists’ attempt to display enzymes involved in the

decomposition of complex polysaccharides, such as cellulose and starch. *S. cerevisiae* could also be used to display phytase, and average ethanol fermentation rate of the resulting whole-cell biocatalyst improved 18% compared to the control strain using grain as raw material [110].

Originally, the major components of cellulosome, including β -glucosidases, endoglucanases cellobiohydrolases, and carboxymethyl cellulase, were displayed on yeast surface independently [111, 112]. To effectively hydrolyze cellulose, these enzymes were co-displayed on yeast surface to form a multi-enzyme catalytic system. The robust yeast display (YSD) system using a-agglutinin receptor as anchoring motif helped to achieve this purpose [113]. Actually, multiple enzyme encoding genes were ligated to *Aga2* gene in one vector and then these enzymes were secreted out of cells to bind on Aga1 resulting in enzyme random co-display [114, 115]. To further control the ratio and exact position on cell wall of the displayed enzymes, the high-affinity interactions between cohesins and dockerins, by which the cellulase components could be held together forming the structural scaffoldin of cellulosome, were employed in YSD system. In this system, a series of cohesins domains were ligated together with the controlled space and fused with Aga2 to bind on yeast cell wall. The displayed enzymes were fused with different dockerins and secreted out of the cell, which then interacted with the cohesin domains in a specific manner [116]. This system allows a series of designed reactions on one whole-cell biocatalyst.

It should be recognized that most of the YSD systems developed for starch and cellulose hydrolysis were based on *S. cerevisiae* as it has the ability to produce ethanol using these carbon sources. Incomplete self-assembly of heterologous minicellulosome on the yeast cell surface was first realized in 2009, and the engineered yeast was capable of hydrolyzing phosphoric acid-swollen cellulose with several times higher ethanol level than that reached by using purified enzymes (1.2–3.5 g/L) [117, 118]. Moreover, yeast surface display of self-assembled minicellulosomes for simultaneous saccharification and fermentation of cellulose to ethanol was reported for the first time in 2010, which generated 1.8 g/L of ethanol [119]. Self-assembly of cellulosomes with two miniscaffoldins on the yeast cell surface was put forwarded in order to increase the production of ethanol based on above researches (Fig. 6). The study used cellulases from mesophilic strain in consideration of thermal compatibility between cellulose hydrolysis and yeast fermentation. The display level of cellulosome increased, and direct fermentation of microcrystalline cellulose to ethanol was realized [120]. Nevertheless, the production of ethanol is still low. The probably reason was excessive metabolic burden on the host and limitation of surface assembly when so many complex and large proteins were simultaneously expressed in cells and presented on the surface. In addition, the self-assembly of cellulosomes needs to be done under the induction of galactose, but the presence of glucose inhibits the use of galactose by yeast cells. So, the assembly process and the fermentation process must be carried out using galactose and cellulose as carbon, respectively, which would decrease cell cellulose saccharification capacity in whole fermentation process.

Applications of Microbial Surface Display in Biosensors

Highly Selective and Sensitive Saccharide Biosensors Based on Oxido-Reductase Surface-Displayed Microorganism

Carbohydrate compounds are not only the main source of energy needed for the maintenance of life activities, but are also important industrial raw materials. Therefore, saccharide analysis

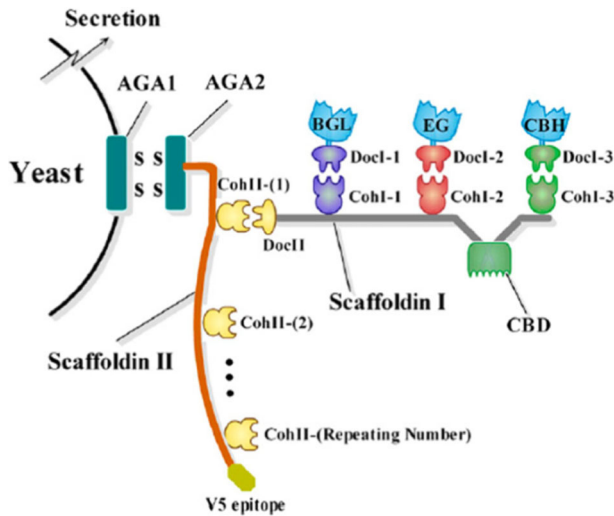


Fig. 6 Cellulosomes with two miniscaffolds self-assembly on the yeast cell surface. Cellulases displayed in this system are as follows: endoglucanase (EG), cellobiohydrolase (exoglucanase) (CBH), and β -glucosidase (BGL). Type I dockerins are as follows: DocI-1, DocI-2, and DocI-3. Type I cohesions are as follows: CohI-1, CohI-2, CohI-3. Type II cohesions are as follows: CohII-(1), CohII-(2), CohII-(repeating number) (Ref. [120])

in food, medicine, and biological processes is extremely important. Traditional detection methods of saccharides suffer from two major drawbacks, i.e., inaccuracy and complicated operation [19]. Several novel biosensors constructed by using enzyme surface-displayed whole cells and nanostructured materials established a rapid and simple method to detect saccharides with high sensitivity and selectivity in our laboratory. Generally, oxido-reductase catalyzed saccharides were displayed on the surfaces of bacteria and yeasts by fusing with anchoring motif, such as INP and α -agglutinin [25, 55]. Multi-walled carbon nanotubes (MWNTs) and oxido-reductase surface displayed cells were cast onto electrode, and concentration of saccharide was determined through measuring the redox signals originating from the above electrode [56]. The resulting biosensor exhibited good stability, wide linear range, and lower detection limit as well as certain anti-interference ability (Fig. 7) [19]. In order to further improve performance of biosensors, rational designs of cell surface-displayed enzymes for increasing stability and selectivity were carried out in our following studies [121]. For example, Nafion/bacteria-displaying glucose dehydrogenase (GDH) mutant/MWNTs composite film-modified electrode was constructed to determine the concentration of glucose [20]. The biosensor prepared with GDH mutant Q252L/E170R/V149K/G259A exhibited good operational stability. Specifically, the amperometric response drifted less than 7% over a period of 1 h, comparing with the wild-type one (10% signal drift during 25 min recording).

Sensitive Electrochemical Microbial Biosensor for Detection of Toxicant

As the most widely used pesticides in the world, organophosphate (OP) insecticides residues are the major causes of food poisoning. A variety of methods have been developed to detect OP pesticides, but there is still no fast, low-cost, and portable approach. As mentioned above, OPH-displayed strain has been exhibiting good detoxification ability. Thus, the whole-cell biocatalyst could be also used to determine the concentration of OP pesticides. In our previous

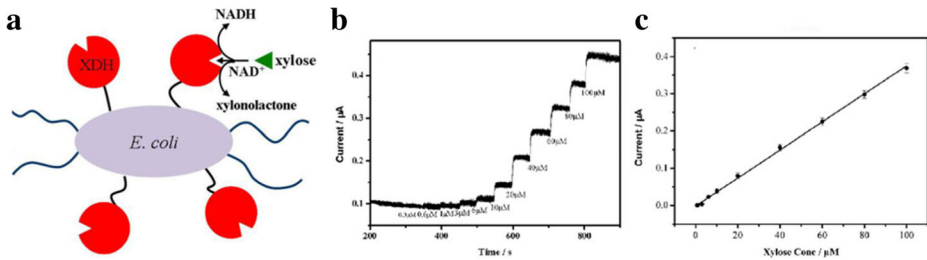


Fig. 7 The construction of a sensitive xylose electrochemical biosensor based on xylose dehydrogenase (XDH) displayed cells and multi-walled carbon nanotubes modified electrode. **(a)** Cell surface-displayed XDH could convert xylose to D-xylonolactone, with NAD^+ as its cofactor. **(b)** Current-time curve obtained for the Nafion/XDH-bacteria/MWNTs/GCE on the successive addition of xylose in 0.1 M PBS (pH 7.4) containing 2 mM NAD^+ , on which the xylose concentration labeled is the xylose concentration in the PBS solution. Applied potential, +0.5 V vs. Ag/AgCl. **(c)** Typical calibration graph of the biosensor for xylose. (Ref. [19])

work, a biosensor based on cell surface-displayed OPH and ordered mesopore carbons for highly specific, sensitive, and rapid detection of *p*-nitrophenyl-substituted OPs compounds was developed [122]. Although, OPH and methyl parathion hydrolase (MPH)-green fluorescent protein (GFP) fusion were co-displayed on the cell surface of *E. coli* using the INP and Lpp-OmpA as the anchoring motifs to further expanded the analyte range in other laboratory, the low detection limit was 100-fold higher than our results [123]. Besides OPH and MPH, acetylcholinesterase (AChE) yeast expressed system could simultaneously detect organic phosphorus and carbamate pesticides with higher sensitivity [124]. However, electrochemical biosensor based on cell surface-displayed AChE was not reported so far. Thus, further study should be focused on the construction of biosensor by integrating AChE-displayed yeast and nanostructured materials, and current efforts in our laboratory are exploring this possibility. In addition to pesticides, toxic heavy metals also threaten safety and security of humans. In order to develop simple, portable, inexpensive, and rapid analytical methods for on-site detecting toxic heavy metals, phytochelatin synthase surface-displayed yeast cell was applied to cadmium biosensors. Unfortunately, the proposed method had room for improvement in terms of sensitivity and selectivity [125].

Application of Microbial Surface Display on Medical Diagnosis

Genetically engineered microbial diagnostic biosensors offer greater sensitivity and convenience compared to classical techniques. Endocrine disrupting compounds (EDCs) are some chemical compounds with similar estrogen activity but different chemical structures, resulting in difficult measurement. To address this issue, Furst et al. developed a novel biosensor based on estrogen receptor alpha ($\text{ER}\alpha$) surface-displayed bacteria, which enabled detections of many detrimental compounds. With this fast and portable device, sub-ppb levels of estradiol and ppm levels of bisphenol A are detected in complex samples. In detail, the novel electrochemical sandwich assay composed of monobodies assembled gold electrode and $\text{ER}\alpha$ surface-displayed bacteria. Changes in the surface impedance could be detected in the presence of EDCs because cells could bind onto EDCs captured gold electrode through the recognition between surfaced-displayed $\text{ER}\alpha$ and EDCs. Compared to free $\text{ER}\alpha$ composed system, the whole cell diagnostic biosensor showed about 100-fold improved sensitivity, which was due to a substantially increased impedance response from the binding of the large cells to the gold surface. Therefore, the small sample size and simple detection system facilitate

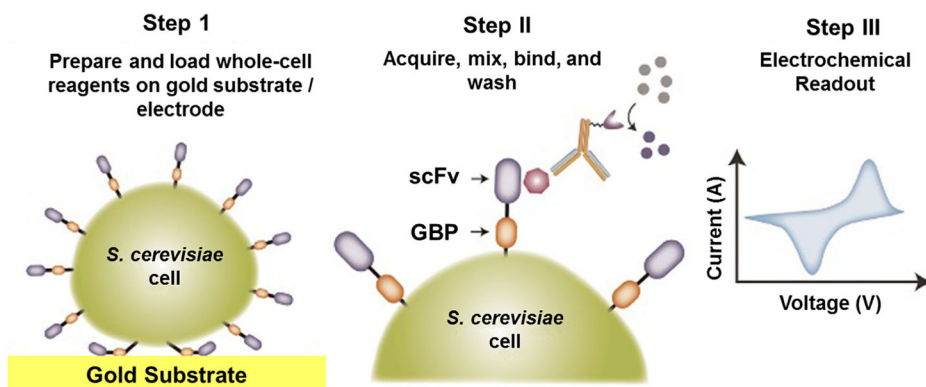


Fig. 8 Schematic illustration of “whole-cell” display of affinity molecules. Step I: After induction, yeast cells are spun and washed and then directly affixed to metal electrodes. Step II: Sample is mixed with detection reagents and applied to the prepared electrode surface and, after incubation, the sample is washed with buffer containing the final detection substrate and redox active molecules. Step III: Detection is carried out on a laboratory or mobile potentiostat (Ref. [127])

its application in the field, and this approach was adequate for measurement of other diverse families of compounds that bind to a single receptor, such as PPAR γ [126]. Similar strategy was also used to detect *Salmonella* outer membrane protein antigen (Fig. 8). Yeast cells were designed to display both single chain variable fragment (scFv) antibodies and gold-binding peptide (GBP) on their surfaces using a-agglutinin as anchoring motif. Sensitive detection of *Salmonella* TM43-E10 surface antigen was realized through sandwich format where the antigen is specifically captured at the cell surface [127]. In order to improve detection sensitivity, the scFv-displayed yeast cell could be fragmented using mortar and pestle with the scFv complex remaining intact to generate nanoyeast-scFvs, which can eliminate interference of larger fragments with electrochemical-based biosensors [128]. Therefore, these studies demonstrated the potential for microbial diagnostic biosensors to serve as robust and low-cost alternatives to current protein-based diagnostic methods.

Conclusion and Prospects

In conclusion, compared with traditional enzymes, surface-displayed enzymes have many advantages. First of all, it is easy to cell culture with high yield and low cost. The copy number of recombinant protein is generally high. Secondly, the enzyme will be inevitably inactivated during the catalytic process or in appropriate environment, whereas the bacterial surface-displayed enzyme can be regenerated through bacterial proliferation. Thirdly, compared with the intracellular enzyme, the cell surface-displayed enzyme exhibits excellent catalytic efficiency and stability. However, the application of this technology is mainly in the laboratory research stage, and the existing surface display system still has shortcomings, such as size limit of passenger and low efficiency of display. Therefore, the development of surface display systems will be the focus of surface display technology research.

As discussed above, carriers in microbial cell surface display system are proteins of outer membrane or cell wall. So, analysis of the proteome of outer membrane or cell wall will be conducive to explore new anchoring motifs. Moreover, artificial display platform could be constructed on the base of well-known carriers, such as spore coat and lipid bilayers. Up to

now, the majority of passengers are displayed on surfaces of bacteria using pathogenic and non-food grade strains as host due to their mature genetic manipulation system, especially *E. coli* [129]. However, this kind of bacteria could hardly be employed in some practical applications, such as in situ degradation of chemical pollutants in the process of food fermentation or antigen delivery vehicle in human body. Therefore, in order to facilitate the presentation of passengers on the surfaces of food-grade, noninvasive, and nonpathogenic bacteria, versatile genetic manipulation and foreign protein expression platforms should be developed through gene engineering means.

Furthermore, the analysis of the tertiary structure of anchoring proteins and in-depth understanding of the role of each component in the transport of passenger protein will help to guide the construction of the cell surface display system and to achieve the functional expression of protein (enzyme). For example, the tertiary structure of several ATs has been analyzed, expounding the role of C-terminal of conserved β -barrel structure in transportation of N-terminal passenger region. However, there are still many unsolved questions about the secretion and folding mechanisms of ATs display system. For example, how is the β -barrel structure of ATs integrated into the outer membrane of the bacteria? The interaction between the passenger region and TU is still lacking in-depth research. Therefore, the further analysis of the tertiary structure of ATs will provide insight into the role of the linked peptide in the interaction between the passenger region and TU, helping to design more reasonable link peptide in the construction of surface display system, realizing the functional display of the proteins (enzymes).

It is worth noting that more kinds of enzymes with different catalytic activities should be displayed on cell surface for various biocatalysis and biosensor. For example, biomass-degradable enzymes could be displayed on surface of *E. coli* or *S. cerevisiae* for production of bioenergy or biochemicals. With the development of biomineralization [130–133] and nanotechnology [134, 135], the integration of cell surface display and versatile inorganic nanomaterials would not only expand scope of the analytes, but also fundamentally improve activity and stability of surface-displayed enzyme as well as sensitivity and stability of biosensor. In addition, so far, present cell surface display-based assays are confined to electrochemical biosensor and spectrophotometric assay. More kinds of analytical methods should be integrated with cell surface display.

In a word, with the development of molecular biology technology and nanotechnology, bacterial and yeast surface display technology will play more and more important role in the fields of biocatalysis and biosensor.

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Compliance with Ethical Standards

Conflict of Interest The authors declare that they have no conflict of interest.

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